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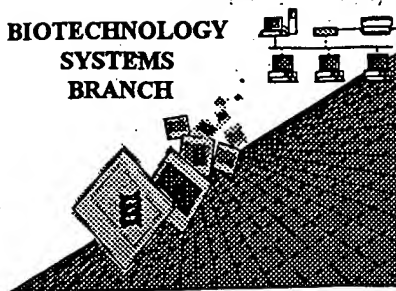
- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ✗ ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
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## **RAW SEQUENCE LISTING** **ERROR REPORT**

BIOTECHNOLOGY  
SYSTEMS  
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/449,817

Source: 1652

Date Processed by STIC: 7/17/2000

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TECH CENTER 1600/2000

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

**FOR FURTHER INFORMATION, PLEASE TELEPHONE MARK SPENCER, 703-308-4212.**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:**

### **Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October-1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

**Checker Version 3.0 can be down loaded from the USPTO website at the following address:**

**<http://www.uspto.gov/web/offices/pac/checker>**

# Raw Sequence Listing Error Summary

## ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/449,817

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 ☐ Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line.  
This may occur if your file was retrieved in a word processor after creating it.  
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 ☐ Wrapped Aminos The amino acid number/text at the end of each line "wrapped" down to the next line.  
This may occur if your file was retrieved in a word processor after creating it.  
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 ☐ Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4 ☐ Misaligned Amino Acid Numbering The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5 ☐ Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.  
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 ☐ Variable Length Sequence(s) contain n's or Xaa's which represented more than one residue.  
As per the rules, each n or Xaa can only represent a single residue.  
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.
- 7 ☐ PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s). Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 8 ☐ Skipped Sequences (OLD RULES) Sequence(s) missing. If intentional, please use the following format for each skipped sequence:  
(2) INFORMATION FOR SEQ ID NO:X:  
(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")  
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:  
This sequence is intentionally skipped  
  
Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 ☐ Skipped Sequences (NEW RULES) Sequence(s) missing. If intentional, please use the following format for each skipped sequence.  
<210> sequence id number  
<400> sequence id number  
000
- 10 ☐ Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.  
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.  
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11 ☒ Use of <213>Organism (NEW RULES) Sequence(s) 1-4 are missing this mandatory field or its response.
- 12 ☐ Use of <220>Feature (NEW RULES) Sequence(s) are missing the <220>Feature and associated headings.  
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"  
Please explain source of genetic material in <220> to <223> section.  
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13 ☐ PatentIn ver. 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).  
Instead, please use "File Manager" or any other means to copy file to floppy disk.

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1652

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/449,817 DATE: 07/17/2000  
 TIME: 11:26:30

Input Set : A:\Seqidt-1.txt  
 Output Set: N:\CRF3\07172000\I449817.raw

Does Not Comply  
 Corrected Diskette Needed

3 <110> APPLICANT: STEINER, MITCHELL; RINALDY, AUGUSTINE; AND MENON, REMA  
 5 <120> TITLE OF INVENTION: AN ISOLATED NUCLEIC ACID ENCODING P-HYDE  
 6 PROTEIN AND METHODS OF INDUCING SUSCEPTIBILITY  
 7 TO INDUCTION OF CELL DEATH IN CANCER  
 9 <130> FILE REFERENCE: P-2762-US1  
 11 <140> CURRENT APPLICATION NUMBER: 09/449,817  
 12 <141> CURRENT FILING DATE: 1999-11-26  
 14 <160> NUMBER OF SEQ ID NOS: 4  
 16 <170> SOFTWARE: PatentIn Ver. 2.1  
 18 <210> SEQ ID NO: 1  
 19 <211> LENGTH: 733  
 20 <212> TYPE: DNA  
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 24 gaggaggagg tctggcggat ggagatctac ctctccctgg gagtgtctgg cctcggcacg 120  
 25 ttgtccctgc tggccgtgac ctcactgccg tccattgcaa actcgtctca ctggaggagg 180  
 26 ttcaagcttc ttcagtcctc actgggcttt gtggccctcg tctgagcac actgcacacg 240  
 27 ctcacctacg gctggaccgg cgccctcgaa gagaccgcta caagttctac ctgcctccca 300  
 28 ccttcacgct cagcgtgctg gtgccctgcg tctcatcct ggccaaagcc ctgtttctcc 360  
 29 tgccttgcac cagccgcaga ctccccagga tccggagaag ctgggagagg gagagcacc 420  
 30 tcaagttcac gctgcccaca gaccacgccc tggccgagaa gacgagccac gtatgagg 480  
 31 cctgccctgg gctctggacc ccgggcacac gagggacggt gccctgagcc cgttagggtt 540  
 32 tottttcttg gtgggtgcaa gtggtataac tgtgtgcaa taggagggtt gaggtccaaa 600  
 33 ttctctggga tcaaatgtat gcatgactat tcagaatgat atacacacat atgtgtatat 660  
 34 gtattttacat atattccaca tatataacag gatttgaat tatacatagc tagctaaaaa 720  
 35 aaaaaaaaaa aaa 733  
 38 <210> SEQ ID NO: 2  
 39 <211> LENGTH: 1467  
 40 <212> TYPE: DNA  
 W--> 42 <213> ORGANISM: *same env*  
 42 <400> SEQUENCE: 2  
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 44 ctggctgagg tccccaagga ggctcccaaa gtgggcatcc tgggcagcgg ggattttgcc 120  
 45 cgggtccctg ccacacgctt ggtgggctct ggcttctttg tgggtggagg aagccgtaac 180  
 46 cccaaacgca ctgcccgcct ctccccctcc tttagcccaag tgactttcca ggaggaggcc 240  
 47 gtgagctctc cagaggatcat ctttgtggcc gtgttccggg agcactactc ctcactgtgc 300  
 48 agtcttgcgt accagttggc tggcaagatc ctatgtgatg taagcaaccc cagggagaag 360  
 49 gagegtcttc agcaccgcca gtccgaacgc gactacctgg cctccctctt cctgcctgc 420  
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 55 cggaaagatg agaacaagtt ctacaagatg cccctgtctg tgggtcaaac cagcataccc 780  
 56 tgtgtggtt acgtgctgct gtccctggtt tacctgctg gtgtgctggc agctgccctt 840  
 57 cagctgagga gggggaccaa gtaccagcgc ttccagact ggctggacca ttggctgcag 900

*see circled  
 portion of  
 Item 12 on  
 Env Summary  
 sheet.*

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/449,817

DATE: 07/17/2000  
TIME: 11:26:30

Input Set : A:\Seqidt-1.txt  
Output Set: N:\CRF3\07172000\I449817.raw

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60 gtccctggcca acaagagccg cctctgggtt gaggaagaag tctggcggtt ggagatatac 1080
61 ctgtccctgg gtgtgctggc tctgggcatg ctgtcactgc tggcggttac ctcgatccct 1140
62 tccattgcaa actcactcaa ctggaaggag ttcagctttg tgcagtccac gctgggcttc 1200
63 gtggccctga tgcctgagcac aatgcacacc ctcacctacg gctggaccgc tgcctttgag 1260
64 gaaaaccact acaagtctta cctgccacc cctcttcttc ctgccctgct cctgccctgt 1320
65 gtcatcatcc tggccaaggg cctcttcttc ctgccctgct tgcaccacag actcaccacg 1380
66 atccgcaggg gctgggagag ggatgggtgc gtcaagttca tgcctgccgc tggccacaca 1440
67 cagggggaga aaacaagcca cgtgtga 1467

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71 <210> SEQ ID NO: 3

72 <211> LENGTH: 3884

73 <212> TYPE: DNA

W--> 75 <213> ORGANISM:

75 <400> SEQUENCE: 3

*same*

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77 ggtggagttt gtgacgtggc gcggggcgctg ggaacggggc ggttgacgta gtagtggtgc 120
78 ggaagtgtga tgttgcaagt gtggcggaac acatgtaagc gacggatgtg gcaaaagtga 180
79 cgtttttggt gtgcgcccgt gtacacagga agtgacaatt ttcgcgcggt tttaggcgga 240
80 tgttgtagta aatttgggcg taaccgagta agattttggc attttcgcg gaaaactgaa 300
81 taagaggag tgaaatctga ataattttgt gttactcata gcgcgtaata tttgtctagg 360
82 gccgcgggga ctttgaccgt ttacgtggag actcgcccag ggcgcgcccc gatgtacggg 420
83 ccagatatac gcgtatctga ggggactagg gtgtgtttag gcgaaaagcg gggcttcggt 480
84 tgtacgcggt taggagttcc ctcaggatat agtagtttcg cttttgcata gggaggggga 540
85 aatgtagtct tatgcaatac tctttagtgc ttgcaacatg gtaacgatga gttagcaaca 600
86 tgccttaaca ggagagaaaa agcaccgtgc atgccgattg gtggaagtaa ggtggtacga 660
87 tctgtccctt tagggaaggg aacagacggg tctgacatgg attggacgaa ccactgaatt 720
88 ccgcattgca gagatattgt atttaagtgc ctgactcgat acaataaacg ccatttgacc 780
89 attcaccaca ttggtgtgca cctccggccc tggccactct cttccgcata gctgtctgcg 840
90 ggggccagct gttgggctcg cggttgagga caaactcttc gcggtcttcc cagtactctt 900
91 ggatcggaac cccgtcgccc tccgaacggt actccgcccg cgagggacct gagcgagtcc 960
92 gcatcgaccg gatcggaaaa cctctcgaga aaggcggtga accagtcaca gtcgctctag 1020
93 aactagtgtg tccccgggct tgcagggaatt cgataattcg gcacgaggct gccgaggcac 1080
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TECH CENTER 1600/2800

## RAW SEQUENCE LISTING

DATE: 07/17/2000

PATENT APPLICATION: US/09/449,817

TIME: 11:26:30

Input Set : A:\Seqidt-1.txt

Output Set: N:\CRF3\07172000\I449817.raw

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117 caagatccgc aggggctggg agagggatgg tgcggtcaag ttcattgtgc ccgctggcca 2520
118 cacacagggg gagaaaacaa gccacgtgtg agggcctgga aatggagaca ggcacagctt 2580
119 gtggggggccc tgggctgggt tgggtctctt tttctgggat ggtatatgcy tgggtggccg 2640
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121 gcgataagta ctcacatata ttcatatat aataggattt actattatc ttagttaaaa 2760
122 aaaaatagtg ggtccttata tttcaactta tgcagggtcc ctatatattc acttgagcat 2820
123 ttcagagcaa atgccacaca ttaaacagca gatcccaccc ttgtggtagc tgcagagaca 2880
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138 aagcttatcg ataccgtcga aacttgttta ttgcagctta taatggttac aaataaagca 3780
139 atagcatcac aaatttcaca aataaagcat ttttttcaat gcattctagt tgtggtttgt 3840
140 ccaaaactcat caatgtatct tatcatgtct ggatccgacc tcg 3884
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143 &lt;210&gt; SEQ ID NO: 4

144 &lt;211&gt; LENGTH: 32166

145 &lt;212&gt; TYPE: DNA

W--&gt; 147 &lt;213&gt; ORGANISM:

147 &lt;400&gt; SEQUENCE: 4

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149 ggtaaacata ttaggaacca gcctgtgatg ctggatgtga ccgaggagct gaggcccgat 120
150 cacttggtgc tggcctgcac ccgcgtgag tttggctcta gcgatgaaga tacagattga 180
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152 atgtagtttt gtatctgttt tgcagcagcc gccgcgcga tgagcaccaa ctcgtttgat 300
153 ggaagcattg tgagctcata ttgacaacg cgcagtcgcc catgggcccgg ggtgcgtcag 360
154 aatgtgatgg gctccagcat tgatggtcgc cccgtcctgc ccgcaaacct tactaccttg 420
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156 gctgcagcca ccgcccgcgg gattgtgact gactttgctt tccagagccc gcttgcaagc 540
157 agtgcagctt cccgttcac gcgccgcgat gacaagttga cggctctttt ggcacaattg 600
158 gattctttga ccgaggaaat taatgtcgtt tctcagcagc tgttggtatc gcgccagcag 660
159 gtttctgccc tgaaggcttc ctcccctccc aatgcggttt aaaacataaa taaaaaacca 720
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## RAW SEQUENCE LISTING

DATE: 07/17/2000

PATENT APPLICATION: US/09/449,817

TIME: 11:26:30

Input Set : A:\Seqidt-1.txt

Output Set: N:\CRF3\07172000\I449817.raw

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166 atgagatgca tcttgactgt tatttttagg ttggtatgt tcccagccat atccctcccg 1140
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169 tccatgcatt cgtccataat gatggcaatg ggcccacggg cgggcgccctg ggccaagata 1320
170 tttctgggat cactaacgtc atagtgtgtt tccaggatga gatcgctata ggccattttt 1380
171 acaaagcgcg ggcggagggt gccagactgc ggtataatgg ttccatcccg cccaggggcg 1440
172 tagttaccct cacagatttg catttccac cctttgagtt cagatggggg gatcatgtct 1500
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174 aggttctctg gcagctgcga cttaccgcag ccggtggggc cgtaaatac acctattacc 1620
175 ggggtcaact ggtagttaa agagctgcag ctgcccgtcat ccctgagcag gggggccact 1680
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177 ccgccacgag atagcagttc ttgcaaggaa gcaaagtttt tcaacgggtt gagaccgtcc 1800
178 gccgtaggca tgcctttgag cgtttgacca agcagttcca ggcgttcca cagctcggtc 1860
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190 gcgtccaggg cagcacgaag gaggctaagt gggaggggta gcggtcgttg tccactaggg 2580
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208 tggttggacg ctggaagacg ttgaagctgg cgtctgtgag acctaccg cg tcacgcacga 3660
209 aggaggcgta ggagtcgcgc agcttgttga ccagctcggc ggtgacctgc acgtctaggg 3720
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## RAW SEQUENCE LISTING

DATE: 07/17/2000

PATENT APPLICATION: US/09/449,817

TIME: 11:26:30

Input Set : A:\Seqidt-1.txt

Output Set: N:\CRF3\07172000\I449817.raw

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/449,817

DATE: 07/17/2000

TIME: 11:26:31

Input Set : A:\Segidt-1.txt

Output Set: N:\CRF3\07172000\I449817.raw

L:22 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:42 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:75 M:282 W: Numeric Field Identifier Missing, <213> is required.  
L:147 M:282 W: Numeric Field Identifier Missing, <213> is required.